

Applicants: Jason D. Bannan and John E. Zabriskie
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Page 2

In the Specification:

Please replace the paragraph beginning on page 13, line 7, with the following rewritten paragraph:

Figure 3. Comparison of the synthetic peptide sequences (SEQ ID NO:3, region 1; and SEQ ID NO:4, region 2) to conserved regions 1 and 2 of the Staphylococcal enterotoxins (SEA (SEQ ID NO:9, region 1; and SEQ ID NO:18, region 2), SEB (SEQ ID NO:10, region 1; and SEQ ID NO:19, region 2), SEC (SEQ ID NO:11, region 1; and SEQ ID NO:20, region 2), SED (SEQ ID NO:12, region 1; and SEQ ID NO:21, region 2), SEE (SEQ ID NO:13, region 1; and SEQ ID NO:22, region 2), and SHE (SEQ ID NO:14, region 1; and SEQ ID NO:23, region 2)), and streptococcal pyrogenic exotoxins (SPEA (SEQ ID NO:15, region 1; and SEQ ID NO:24, region 2), SPEC (SEQ ID NO:16, region 1; and SEQ ID NO:25, region 2) and SSA (SEQ ID NO:17, region 1; and SEQ ID NO:26, region 2)). Staphylococcal toxic shock syndrome toxin 1 (TSST-1, SEQ ID NO:27) was compared with the region 2 peptide. Numbers represent the residue positions as a reference to where these regions exist in the whole toxin molecules. Sequences are from either the Swiss protein or GenBank databases under the following accession numbers. Swiss protein: SPEA, P08095; SPEC, P13380; SEA, P13163; SEB, P01552; SEC, P01553; SED, P20723; SEE, P12993. GenBank: SEH, U11702; SSA, L29565; TSST1, J02615.